PCT10

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/089,986

DATE: 04/25/2002
TIME: 11:51:22

Input Set : A:\EP.txt
Output Set: N:\CRF3\04252002\J089986.raw

```
      3 <110> APPLICANT: KUROSAWA, Keiko
      4       KAJIYAMA, Naoki
      6 <120> TITLE OF INVENTION: GENE ENCODING PROTEIN HAVING THE ABILITY TO REGENERATE
LUCIFERIN, NOVEL
      7       RECOMBINANT DNA, PROCESS FOR THE PREPARATION OF PROTEIN HAVING THE ABILITY TO
      8       REGENERATE LUCIFERIN
     10 <130> FILE REFERENCE: 221609US0PCT
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/089,986
C--> 12 <141> CURRENT FILING DATE: 2002-04-08
     12 <150> PRIOR APPLICATION NUMBER: PCT/JP00/06527
     13 <151> PRIOR FILING DATE: 2000-09-22
     15 <150> PRIOR APPLICATION NUMBER: JP285258/1999
     16 <151> PRIOR FILING DATE: 1999-10-06
     18 <160> NUMBER OF SEQ ID NOS: 11
     20 <170> SOFTWARE: PatentIn version 3.1
     22 <210> SEQ ID NO: 1
     23 <211> LENGTH: 924
     24 <212> TYPE: DNA
     25 <213> ORGANISM: Photinus pyralis
     27 <220> FEATURE:
     28 <221> NAME/KEY: misc_feature
     29 <222> LOCATION: (843)..(843)
     30 <223> OTHER INFORMATION: n is a, g, t, or c
     33 <400> SEQUENCE: 1
     34 atggggccag ttgttgaaaa aattgcagaa cttggcaagt atacggttgg agaaggtcct     60
     36 cactgggatc atgaaactca gaccttatat ttcgtcgaca ccgtagagaa aacttttcat    120
     38 aaatatgtac cttctcagaa aaaatacacg ttttgtaaag tagataaact ggtttctttc    180
     40 attattcccc ttgctggatc ccctggccgt tttgtagtca gtttggaacg tgaaatagcc    240
     42 attcttacat gggatggcgt tagtgctgca cctacaagca tagaagctat tgttaatgtc    300
     44 gaaccacaca ttaaaaataa cagactcaat gatggcaaag cagatcccct tggcaatcta    360
     46 tggacaggta caatggctat tgacgctggt ctccccgtag gaccggtcac tggcagttta    420
     48 tatcatttag gggctgataa aaaggtaaaa atgcacgaga gcaacatagc tatagcaaat    480
     50 gggctcgcgt ggagtaatga tttgaagaaa atgtattata ttgattcggg gaaaagaaga    540
     52 gtagacgagt acgattatga tgcttctaca ttatccatca gcaatcaacg gccattattt    600
     54 acttttgaaa agcatgaagt gcctggatat ccagatggtc aaacaattga tgaggagggt    660
     56 aatttatggg ttgccgtttt ccaaggacag cgaattatta aaatcagtac ccaacaaccg    720
     58 gaagtgttac tggataccgt aaaaatacca gatcctcagg tcacctctgt agcatttggc    780
     60 ggtccgaatt tggatgaact gcatgtaaca tctgctggtc ttcagcttga cgacagttct    840
W--> 62 ttngacaaaa gtttagttaa tgggcacgtc tacagagtaa caggtttagg cgtcaaaggt    900
     62                                                                    924
     64 ttcgcgggag ttaaagtgaa gcta
     67 <210> SEQ ID NO: 2
     68 <211> LENGTH: 308
     69 <212> TYPE: PRT
     70 <213> ORGANISM: Photinus pyralis
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<400> SEQUENCE: 2

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Met Gly Pro Val Val Glu Lys Ile Ala Glu Leu Gly Lys Tyr Thr Val
1               5                  10                  15
Gly Glu Gly Pro His Trp Asp His Glu Thr Gln Thr Leu Tyr Phe Val
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Asp Thr Val Glu Lys Thr Phe His Lys Tyr Val Pro Ser Gln Lys Lys
        35                  40                  45
Tyr Thr Phe Cys Lys Val Asp Lys Leu Val Ser Phe Ile Ile Pro Leu
    50                  55                  60
Ala Gly Ser Pro Gly Arg Phe Val Val Ser Leu Glu Arg Glu Ile Ala
65                  70                  75                  80
Ile Leu Thr Trp Asp Gly Val Ser Ala Ala Pro Thr Ser Ile Glu Ala
                85                  90                  95
Ile Val Asn Val Glu Pro His Ile Lys Asn Asn Arg Leu Asn Asp Gly
            100                 105                 110
Lys Ala Asp Pro Leu Gly Asn Leu Trp Thr Gly Thr Met Ala Ile Asp
        115                 120                 125
Ala Gly Leu Pro Val Gly Pro Val Thr Gly Ser Leu Tyr His Leu Gly
    130                 135                 140
Ala Asp Lys Lys Val Lys Met His Glu Ser Asn Ile Ala Ile Ala Asn
145                 150                 155                 160
Gly Leu Ala Trp Ser Asn Asp Leu Lys Lys Met Tyr Tyr Ile Asp Ser
                165                 170                 175
Gly Lys Arg Arg Val Asp Glu Tyr Asp Tyr Asp Ala Ser Thr Leu Ser
            180                 185                 190
Ile Ser Asn Gln Arg Pro Leu Phe Thr Phe Glu Lys His Glu Val Pro
        195                 200                 205
Gly Tyr Pro Asp Gly Gln Thr Ile Asp Glu Glu Gly Asn Leu Trp Val
    210                 215                 220
Ala Val Phe Gln Gly Gln Arg Ile Ile Lys Ile Ser Thr Gln Gln Pro
225                 230                 235                 240
Glu Val Leu Leu Asp Thr Val Lys Ile Pro Asp Pro Gln Val Thr Ser
                245                 250                 255
Val Ala Phe Gly Gly Pro Asn Leu Asp Glu Leu His Val Thr Ser Ala
            260                 265                 270
Gly Leu Gln Leu Asp Asp Ser Ser Leu Asp Lys Ser Leu Val Asn Gly
        275                 280                 285
His Val Tyr Arg Val Thr Gly Leu Gly Val Lys Gly Phe Ala Gly Val
    290                 295                 300
Lys Val Lys Leu
305
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<210> SEQ ID NO: 3
<211> LENGTH: 24
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: synthetic DNA
<400> SEQUENCE: 3
gttggagaag gaccgatttg ggat 24

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Input Set : A:\EP.txt
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166 <210> SEQ ID NO: 4
167 <211> LENGTH: 29
168 <212> TYPE: DNA
169 <213> ORGANISM: Artificial Sequence
171 <220> FEATURE:
172 <223> OTHER INFORMATION: synthetic DNA
174 <400> SEQUENCE: 4 29
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178 <210> SEQ ID NO: 5
179 <211> LENGTH: 20
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: synthetic DNA
186 <400> SEQUENCE: 5 20
187 ggacaggtac aatggctatt
190 <210> SEQ ID NO: 6
191 <211> LENGTH: 20
192 <212> TYPE: DNA
193 <213> ORGANISM: Artificial Sequence
195 <220> FEATURE:
196 <223> OTHER INFORMATION: synthetic DNA
198 <400> SEQUENCE: 6 20
199 atcgtactcg tctactcttc
202 <210> SEQ ID NO: 7
203 <211> LENGTH: 22
204 <212> TYPE: DNA
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: synthetic DNA
210 <400> SEQUENCE: 7 22
211 taggtgcagc actaacgcca tc
214 <210> SEQ ID NO: 8
215 <211> LENGTH: 21
216 <212> TYPE: DNA
217 <213> ORGANISM: Artificial Sequence
219 <220> FEATURE:
220 <223> OTHER INFORMATION: synthetic DNA
222 <400> SEQUENCE: 8 21
223 ttcacgttcc aaactgacta c
226 <210> SEQ ID NO: 9
227 <211> LENGTH: 23
228 <212> TYPE: DNA
229 <213> ORGANISM: Artificial Sequence
231 <220> FEATURE:
232 <223> OTHER INFORMATION: synthetic DNA
234 <400> SEQUENCE: 9 23
235 ctcgcgtgga gtaatgattt gaa
238 <210> SEQ ID NO: 10

RAW SEQUENCE LISTING
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Input Set : A:\EP.txt
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239 <211> LENGTH: 33
240 <212> TYPE: DNA
241 <213> ORGANISM: Artificial Sequence
243 <220> FEATURE:
244 <223> OTHER INFORMATION: synthetic DNA
246 <400> SEQUENCE: 10 33
247 ggaattcatg gggccagttg ttgaaaaaat tgc
250 <210> SEQ ID NO: 11
251 <211> LENGTH: 35
252 <212> TYPE: DNA
253 <213> ORGANISM: Artificial Sequence
255 <220> FEATURE:
256 <223> OTHER INFORMATION: synthetic DNA
258 <400> SEQUENCE: 11 35
259 aactgcagtc atagcttcac tttaactccc gcgaa

RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 843

VERIFICATION SUMMARY
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L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:62 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:840